

# Resource Summary Report

Generated by [dkNET](#) on Aug 25, 2026

## pcDNA3.1 GW-NL-myc

RRID:Addgene\_113447

Type: Plasmid

### Proper Citation

RRID:Addgene\_113447

### Plasmid Information

**URL:** <http://www.addgene.org/113447>

**Proper Citation:** RRID:Addgene\_113447

**Bacterial Resistance:** Chloramphenicol and Ampicillin

**Defining Citation:** [PMID:29997244](https://pubmed.ncbi.nlm.nih.gov/29997244/)

**Vector Backbone Description:** Backbone Size:5428; Vector Backbone:pcDNA3.1; Vector Types:Mammalian Expression, Luciferase, Gateway shuttle vector; Bacterial Resistance:Chloramphenicol and Ampicillin

**Plasmid Name:** pcDNA3.1 GW-NL-myc

**Record Creation Time:** 20220422T221554+0000

**Record Last Update:** 20260815T080212+0000

### Ratings and Alerts

No rating or validation information has been found for pcDNA3.1 GW-NL-myc.

No alerts have been found for pcDNA3.1 GW-NL-myc.

### Data and Source Information

**Source:** [Addgene](#)

### Usage and Citation Metrics

We found 5 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [dkNET](#) .

Trepte P, et al. (2024) AI-guided pipeline for protein-protein interaction drug discovery identifies a SARS-CoV-2 inhibitor. *Molecular systems biology*, 20(4), 428.

Lee CY, et al. (2024) Systematic discovery of protein interaction interfaces using AlphaFold and experimental validation. *Molecular systems biology*, 20(2), 75.

Brenner D, et al. (2024) A TBK1 variant causes autophagolysosomal and motoneuron pathology without neuroinflammation in mice. *The Journal of experimental medicine*, 221(5).

Trepte P, et al. (2023) AI-guided pipeline for protein-protein interaction drug discovery identifies a SARS-CoV-2 inhibitor. *bioRxiv : the preprint server for biology*.

Greco TM, et al. (2022) Dynamics of huntingtin protein interactions in the striatum identifies candidate modifiers of Huntington disease. *Cell systems*, 13(4), 304.